

AMENDMENTS TO THE SPECIFICATION:

Please replace the paragraph identified as item 3 on page 24 with the following amended paragraph:

- 3) Standard methods were used for collecting genes from the patients, identifying genes, and specifying diverse amino acids and base sequences. (MCH & IRS, Supplement Vol.1 73-95, 1994. Tissue Antigens 39:187-202, 1992. 38 :53-59,1991, 38 :60-71,1991, 40 :100-103,1992, all of which are incorporated herein by reference in their entireties). For amplification of genomic DNA, for example, an automated PCR thermal sequencer can be used, as described in Tissue Antigens, 38: 53-59, 1991. Analyses positions were from -29 to 237th on DRB1*, -32 to 237th on DQB1*, and -29 to 229th on DPB1*.